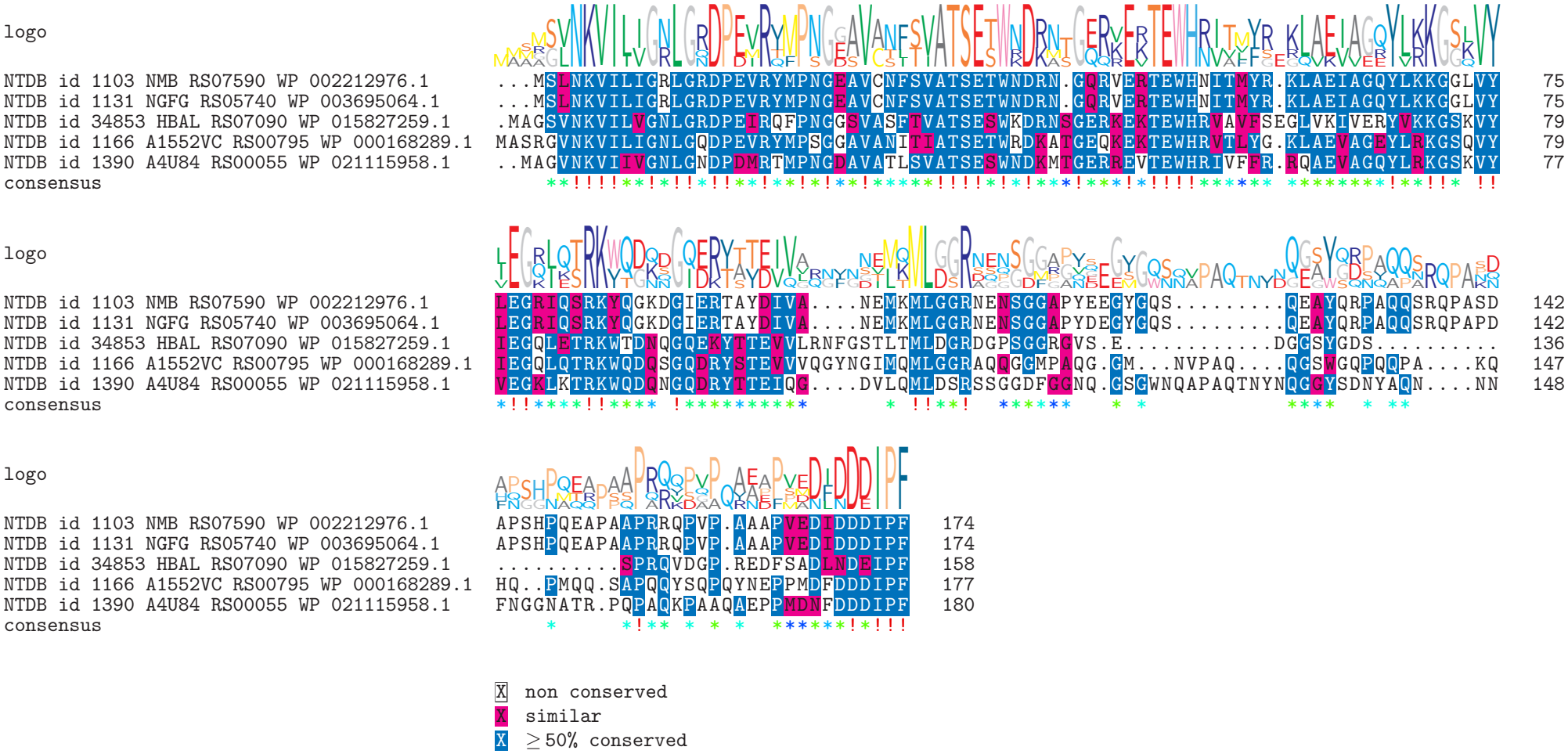
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY 75
NTDB id 1131 NGFG RS05740 WP 003695064.1 ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY 75
NTDB id 34853 HBAL RS07090 WP 015827259.1 .MAGSVNKVILVGNLGRDPEIRQFPNGGSVASFTVATSESWKDRNSGERKEKTEWHRVAVFSEGLVKIVERYVKKGSKVY 79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY 79
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY 77
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQS.........QEAYQRPAQQSRQPASD 142
NTDB id 1131 NGFG RS05740 WP 003695064.1 LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQS.........QEAYQRPAQQSRQPAPD 142
NTDB id 34853 HBAL RS07090 WP 015827259.1 IEGQLETRKWTDNQGQEKYTTEVVLRNFGSTLTMLDGRDGPSGGRGVS.E............DGGSYGDS.......... 136
NTDB id 1166 A1552VC RS00795 WP 000168289.1 IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQG.GM...NVPAQ....QGSWGQPQQPA....KQ 147
NTDB id 1390 A4U84 RS00055 WP 021115958.1 VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQ.GSGWNQAPAQTNYNQGGYSDNYAQN....NN 148
consensus
logo
NTDB id 1103 NMB RS07590 WP 002212976.1 APSHPQEAPAAPRRQPVP.AAAPVEDIDDDIPF 174
NTDB id 1131 NGFG RS05740 WP 003695064.1 APSHPQEAPAAPRRQPVP.AAAPVEDIDDDIPF 174
NTDB id 34853 HBAL RS07090 WP 015827259.1 ..........SPRQVDGP.REDFSADLNDEIPF 158
NTDB id 1166 A1552VC RS00795 WP 000168289.1 HQ..PMQQ.SAPQQYSQPQYNEPPMDFDDDIPF 177
NTDB id 1390 A4U84 RS00055 WP 021115958.1 FNGGNATR.PQPAQKPAAQAEPPMDNFDDDIPF 180
consensus
X non conserved
X similar
X ≥ 50% conserved